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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/040,906A

DATE: 07/17/2002

TIME: 13:48:32

Input Set : N:\paola\10040906.raw

Output Set: N:\CRF3\07172002\J040906A.raw

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1 <110> APPLICANT: Arnaut, Greta
2   Boets, Annemie
3   Vanneste, Stijn
4   Van Rie, Jeroen
5   Van Houdt, Sara
6 <120> TITLE OF INVENTION: Novel Bacillus thuringiensis insecticidal proteins
7 <130> FILE REFERENCE: 58764.000036
8 <140> CURRENT APPLICATION NUMBER: US/10/040,906A
9 <141> CURRENT FILING DATE: 2002-06-24
10 <160> NUMBER OF SEQ ID NOS: 9
11 <170> SOFTWARE: PatentIn version 3.1
13 <210> SEQ ID NO: 1
14 <211> LENGTH: 1899
15 <212> TYPE: DNA
16 <213> ORGANISM: Bacillus thuringiensis
17 <220> FEATURE:
18 <221> NAME/KEY: CDS
19 <222> LOCATION: (1)..(1896)
20 <223> OTHER INFORMATION:
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25   aat gta gtg gcc cat gat cca ttt agt ttt gag cat aaa tca tta gat      96
26   Asn Val Val Ala His Asp Pro Phe Ser Phe Glu His Lys Ser Leu Asp
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28   acc atc cga aaa gaa tgg atg gag tgg aaa aga aca gat cat agt tta      144
29   Thr Ile Arg Lys Glu Trp Met Glu Trp Lys Arg Thr Asp His Ser Leu
30   35          40          45
31   tat gta gct cct ata gtc gga act gtt tct agc ttt ctg cta aag aag      192
32   Tyr Val Ala Pro Ile Val Gly Thr Val Ser Ser Phe Leu Leu Lys Lys
33   50          55          60
34   gtg ggg agt ctt att gga aaa agg ata ttg agt gaa tta tgg ggg tta      240
35   Val Gly Ser Leu Ile Gly Lys Arg Ile Leu Ser Glu Leu Trp Gly Leu
36   65          70          75          80
37   ata ttt cct agt ggt agc aca aat cta atg caa gat att tta agg gag      288
38   Ile Phe Pro Ser Gly Ser Thr Asn Leu Met Gln Asp Ile Leu Arg Glu
39   85          90          95
40   aca gaa caa ttc cta aat caa aga ctt aat aca gac act ctt gcc cgt      336
41   Thr Glu Gln Phe Leu Asn Gln Arg Leu Asn Thr Asp Thr Leu Ala Arg
42   100         105         110
43   gta aat gcg gaa ttg gaa ggg ctg caa gcg aat ata agg gag ttt aat      384
44   Val Asn Ala Glu Leu Glu Gly Leu Gln Ala Asn Ile Arg Glu Phe Asn

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95      Gly Val Asn Thr Val Thr Asn Trp Gln Thr Glu Ser Phe Glu Ser Thr
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97      tta ggt tta agg tgt ggt gct ttt aca gct cgt ggt aat tca aac tat      1248
98      Leu Gly Leu Arg Cys Gly Ala Phe Thr Ala Arg Gly Asn Ser Asn Tyr
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100     ttc cca gat tat ttt atc cgt aat att tca gga gtt cct tta gtt gtt      1296
101     Phe Pro Asp Tyr Phe Ile Arg Asn Ile Ser Gly Val Pro Leu Val Val
102     420                               425                               430
103     aga aat gaa gat tta aga aga ccg tta cac tat aat gaa ata aga aat      1344
104     Arg Asn Glu Asp Leu Arg Arg Pro Leu His Tyr Asn Glu Ile Arg Asn
105     435                               440                               445
106     ata gaa agt cct tca gga aca cct ggt gga tta cga gct tat atg gta      1392
107     Ile Glu Ser Pro Ser Gly Thr Pro Gly Gly Leu Arg Ala Tyr Met Val
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109     tct gtg cat aat aga aaa aat aat atc tat gcc gtg cat gaa aat ggt      1440
110     Ser Val His Asn Arg Lys Asn Asn Ile Tyr Ala Val His Glu Asn Gly
111     465                               470                               475                               480
112     act atg att cat tta gcg ccg gaa gat tat aca gga ttc acc ata tcg      1488
113     Thr Met Ile His Leu Ala Pro Glu Asp Tyr Thr Gly Phe Thr Ile Ser
114     485                               490                               495
115     ccg ata cat gca act caa gtg aat aat caa acg cga aca ttt att tct      1536
116     Pro Ile His Ala Thr Gln Val Asn Asn Gln Thr Arg Thr Phe Ile Ser
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118     gaa aaa ttt gga aat caa ggt gat tcc tta aga ttt gaa caa agc aac      1584
119     Glu Lys Phe Gly Asn Gln Gly Asp Ser Leu Arg Phe Glu Gln Ser Asn
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124     tat tta aga gta tct tca cta gga aat tcc act att cga gtt act ata      1680
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130     gat gga gtt aat gat aat ggc gct cgt ttt tta gat att aat atg ggt      1776
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137     Thr Phe Asn Ser Gly Thr Gln Phe Glu Leu Met Asn Ile Met Phe Val
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139     cca act aat ctt cca cca ata tat taa      1899
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141     625                               630
143 <210> SEQ ID NO: 2

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144 <211> LENGTH: 632
145 <212> TYPE: PRT
146 <213> ORGANISM: Bacillus thuringiensis
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152 Thr Ile Arg Lys Glu Trp Met Glu Trp Lys Arg Thr Asp His Ser Leu
153 35 40 45
154 Tyr Val Ala Pro Ile Val Gly Thr Val Ser Ser Phe Leu Leu Lys Lys
155 50 55 60
156 Val Gly Ser Leu Ile Gly Lys Arg Ile Leu Ser Glu Leu Trp Gly Leu
157 65 70 75 80
158 Ile Phe Pro Ser Gly Ser Thr Asn Leu Met Gln Asp Ile Leu Arg Glu
159 85 90 95
160 Thr Glu Gln Phe Leu Asn Gln Arg Leu Asn Thr Asp Thr Leu Ala Arg
161 100 105 110
162 Val Asn Ala Glu Leu Glu Gly Leu Gln Ala Asn Ile Arg Glu Phe Asn
163 115 120 125
164 Gln Gln Val Asp Asn Phe Leu Asn Pro Thr Gln Asn Pro Val Pro Leu
165 130 135 140
166 Ser Ile Thr Ser Ser Val Asn Thr Met Gln Gln Leu Phe Leu Asn Arg
167 145 150 155 160
168 Leu Pro Gln Phe Arg Val Gln Gly Tyr Gln Leu Leu Leu Leu Pro Leu
169 165 170 175
170 Phe Ala Gln Ala Ala Asn Met His Leu Ser Phe Ile Arg Asp Val Val
171 180 185 190
172 Leu Asn Ala Asp Glu Trp Gly Ile Ser Ala Ala Thr Leu Arg Thr Tyr
173 195 200 205
174 Gln Asn Tyr Leu Lys Asn Tyr Thr Thr Glu Tyr Ser Asn Tyr Cys Ile
175 210 215 220
176 Asn Thr Tyr Gln Thr Ala Phe Arg Gly Leu Asn Thr Arg Leu His Asp
177 225 230 235 240
178 Met Leu Glu Phe Arg Thr Tyr Met Phe Leu Asn Val Phe Glu Tyr Val
179 245 250 255
180 Ser Ile Trp Ser Leu Phe Lys Tyr Gln Ser Leu Leu Val Ser Ser Gly
181 260 265 270
182 Ala Asn Leu Tyr Ala Ser Gly Ser Gly Pro Gln Gln Thr Gln Ser Phe
183 275 280 285
184 Thr Ser Gln Asp Trp Pro Phe Leu Tyr Ser Leu Phe Gln Val Asn Ser
185 290 295 300
186 Asn Tyr Val Leu Asn Gly Phe Ser Gly Ala Arg Leu Thr Gln Thr Phe
187 305 310 315 320
188 Pro Asn Ile Gly Gly Leu Pro Gly Thr Thr Thr Thr His Ala Leu Leu
189 325 330 335
190 Ala Ala Arg Val Asn Tyr Ser Gly Gly Val Ser Ser Gly Asp Ile Gly
191 340 345 350
192 Ala Val Phe Asn Gln Asn Phe Ser Cys Ser Thr Phe Leu Pro Pro Leu

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194    Leu Thr Pro Phe Val Arg Ser Trp Leu Asp Ser Gly Ser Asp Arg Gly
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197          385          390          395          400
198    Leu Gly Leu Arg Cys Gly Ala Phe Thr Ala Arg Gly Asn Ser Asn Tyr
199          405          410          415
200    Phe Pro Asp Tyr Phe Ile Arg Asn Ile Ser Gly Val Pro Leu Val Val
201          420          425          430
202    Arg Asn Glu Asp Leu Arg Arg Pro Leu His Tyr Asn Glu Ile Arg Asn
203          435          440          445
204    Ile Glu Ser Pro Ser Gly Thr Pro Gly Gly Leu Arg Ala Tyr Met Val
205          450          455          460
206    Ser Val His Asn Arg Lys Asn Asn Ile Tyr Ala Val His Glu Asn Gly
207          465          470          475          480
208    Thr Met Ile His Leu Ala Pro Glu Asp Tyr Thr Gly Phe Thr Ile Ser
209          485          490          495
210    Pro Ile His Ala Thr Gln Val Asn Asn Gln Thr Arg Thr Phe Ile Ser
211          500          505          510
212    Glu Lys Phe Gly Asn Gln Gly Asp Ser Leu Arg Phe Glu Gln Ser Asn
213          515          520          525
214    Thr Thr Ala Arg Tyr Thr Leu Arg Gly Asn Gly Asn Ser Tyr Asn Leu
215          530          535          540
216    Tyr Leu Arg Val Ser Ser Leu Gly Asn Ser Thr Ile Arg Val Thr Ile
217          545          550          555          560
218    Asn Gly Arg Val Tyr Thr Ala Ser Asn Val Asn Thr Thr Thr Asn Asn
219          565          570          575
220    Asp Gly Val Asn Asp Asn Gly Ala Arg Phe Leu Asp Ile Asn Met Gly
221          580          585          590
222    Asn Val Val Ala Ser Asp Asn Thr Asn Val Pro Leu Asp Ile Asn Val
223          595          600          605
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225          610          615          620
226    Pro Thr Asn Leu Pro Pro Ile Tyr
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229 <210> SEQ ID NO: 3
230 <211> LENGTH: 1899
231 <212> TYPE: DNA
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234 <221> NAME/KEY: CDS
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236 <223> OTHER INFORMATION:
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239    Met Asn Ser Val Leu Asn Ser Gly Arg Thr Thr Ile Cys Asp Ala Tyr
240    1          5          10          15
241    aat gta gtg gct cat gat cca ttt agt ttt caa cat aaa tca tta gat      96
242    Asn Val Val Ala His Asp Pro Phe Ser Phe Gln His Lys Ser Leu Asp

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VERIFICATION SUMMARY

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